

**AMENDMENTS TO THE SPECIFICATION**

Please amend the paragraph beginning at page 5, line 22 as follows:

Figures 2A-2C show a semi-quantitative RT-PCR analysis of 20P2H8 gene expression in a panel of 16 normal human tissues (Panels 2B and 2C) and several prostate cancer xenografts (Panel 2A). Lanes 1-8 in panel 2A correspond to RT-PCR analysis of 20P2H8 gene expression in brain, prostate, LAPC-4 AD, LAPC-4 AI, LAPC-9 AD, HeLa, mouse mix and a negative control respectively. Lanes 1-8 in panel 2B correspond to RT-PCR analysis of 20P2H8 gene expression in brain, heart, kidney, liver, lung, pancreas, placenta and skeletal muscle respectively. Lanes 1-8 in panel 2C correspond to RT-PCR analysis of 20P2H8 gene expression in colon, ovary, leukocytes, prostate, small intestine, spleen, testis and thymus respectively. The RT-PCR primers are specific for 20P2H8 and are as follows: 5'-TCTTGAAACCTCCAGACACAAGAA-3' and 5'-AAGTTACGATTTGGCTTCACTGG-3' (SEQ ID NOS:13 and 16, respectively). The 20P2H8 PCR product is expected to be of 162 base pairs as marked by an arrow.